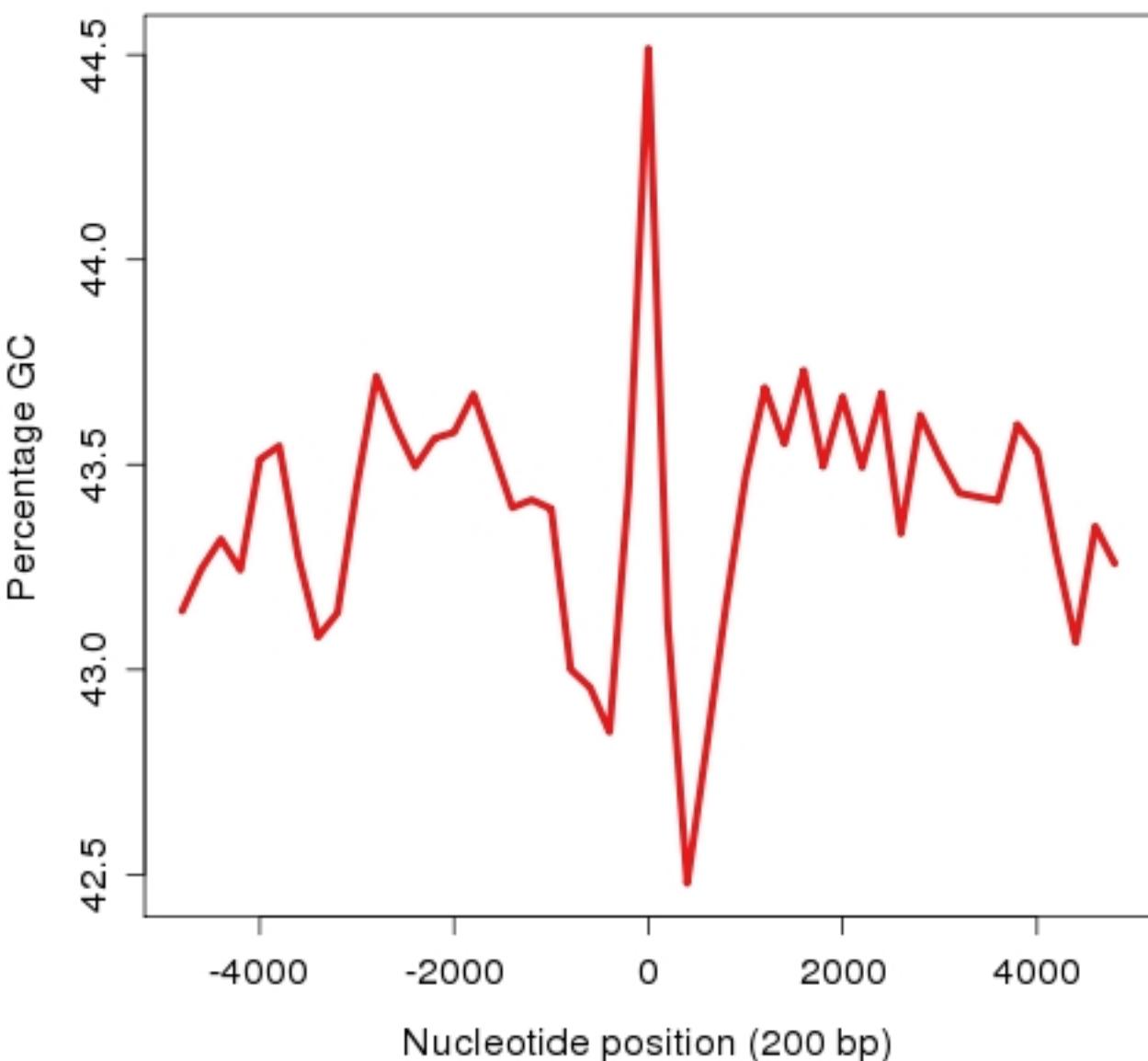
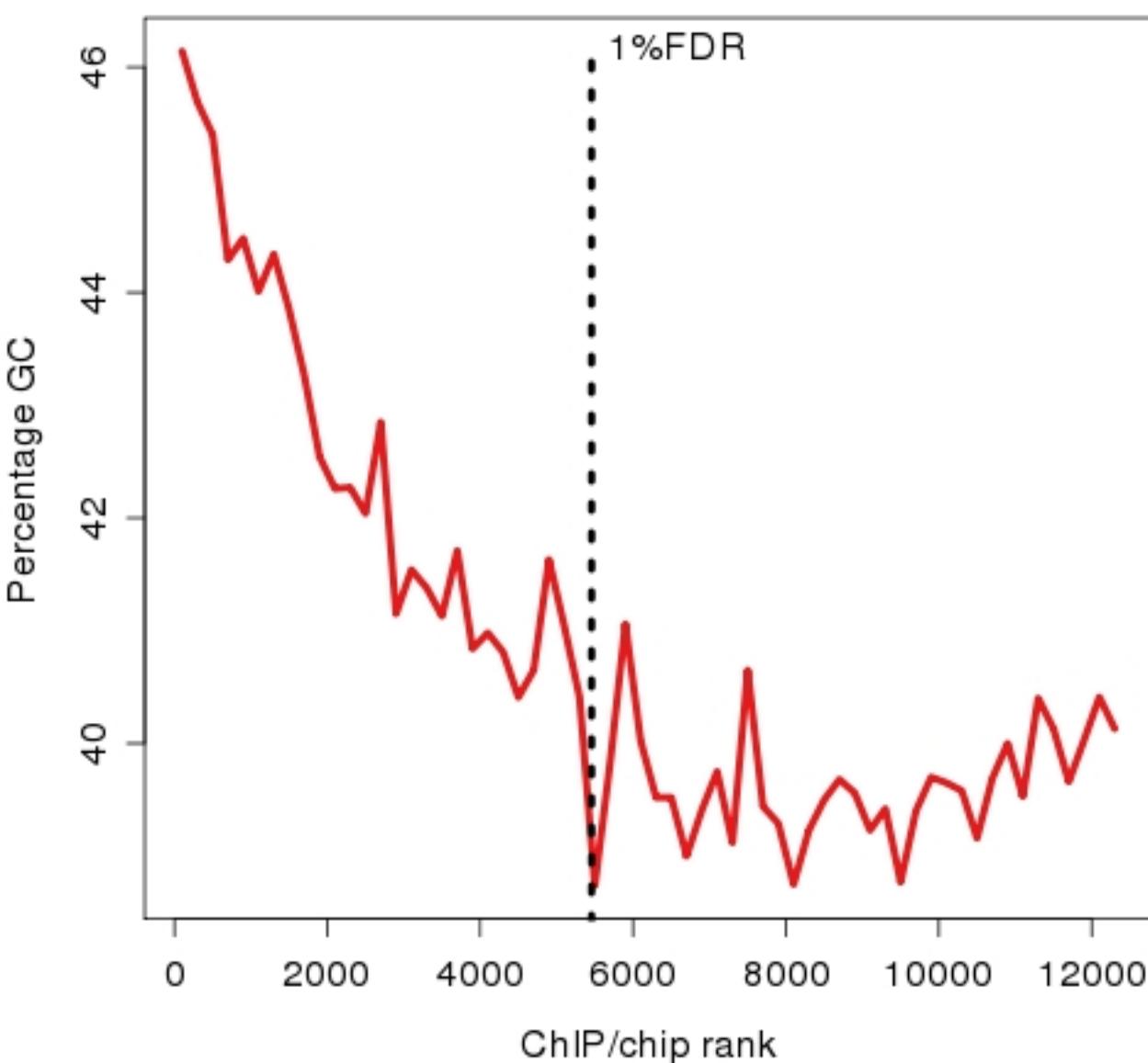


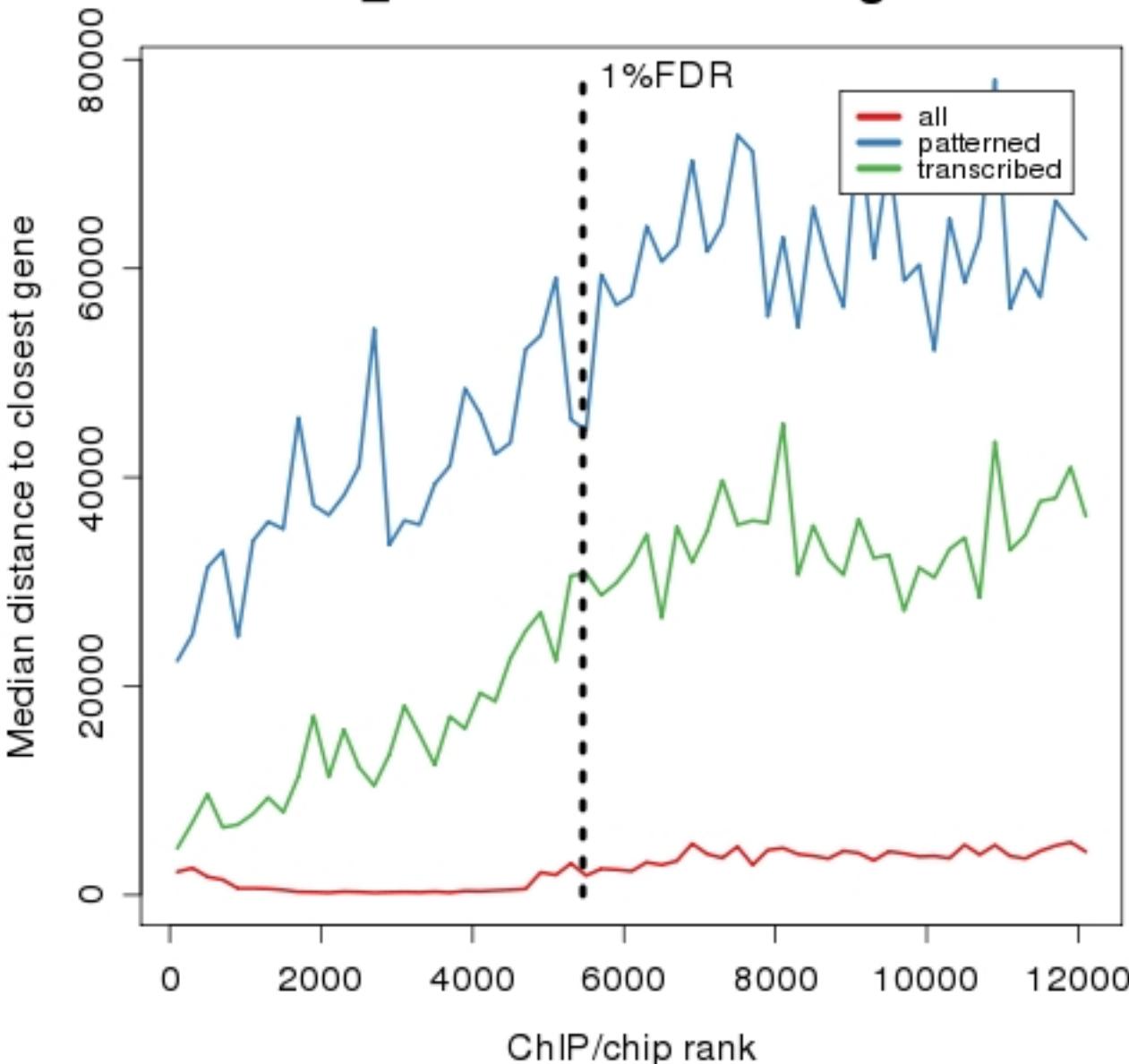
med_2 base composition across peaks



med_2 base composition down ranks



med_2 median distance to genes



med_2 distribution of 1%FDR peak scores

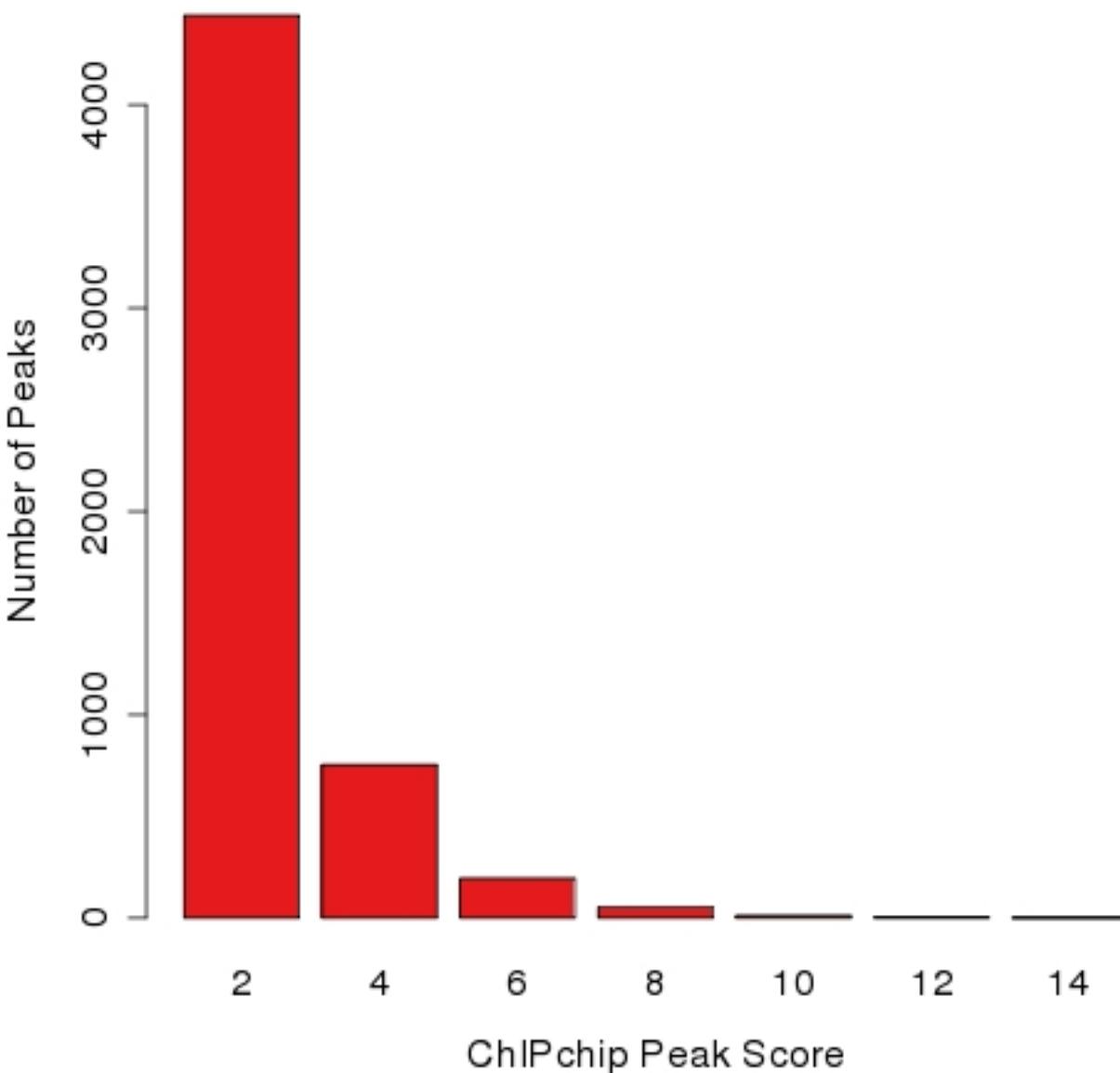
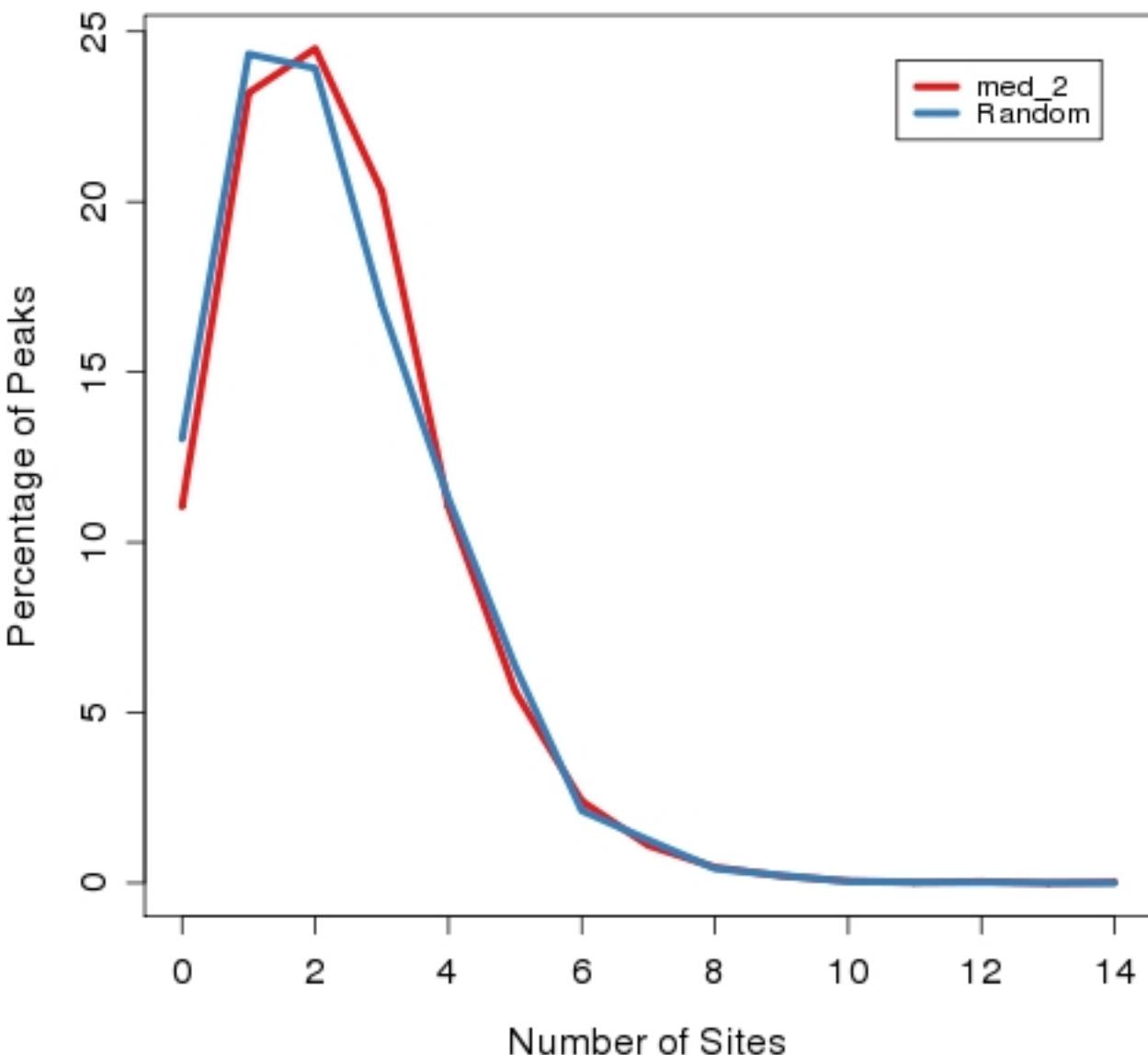
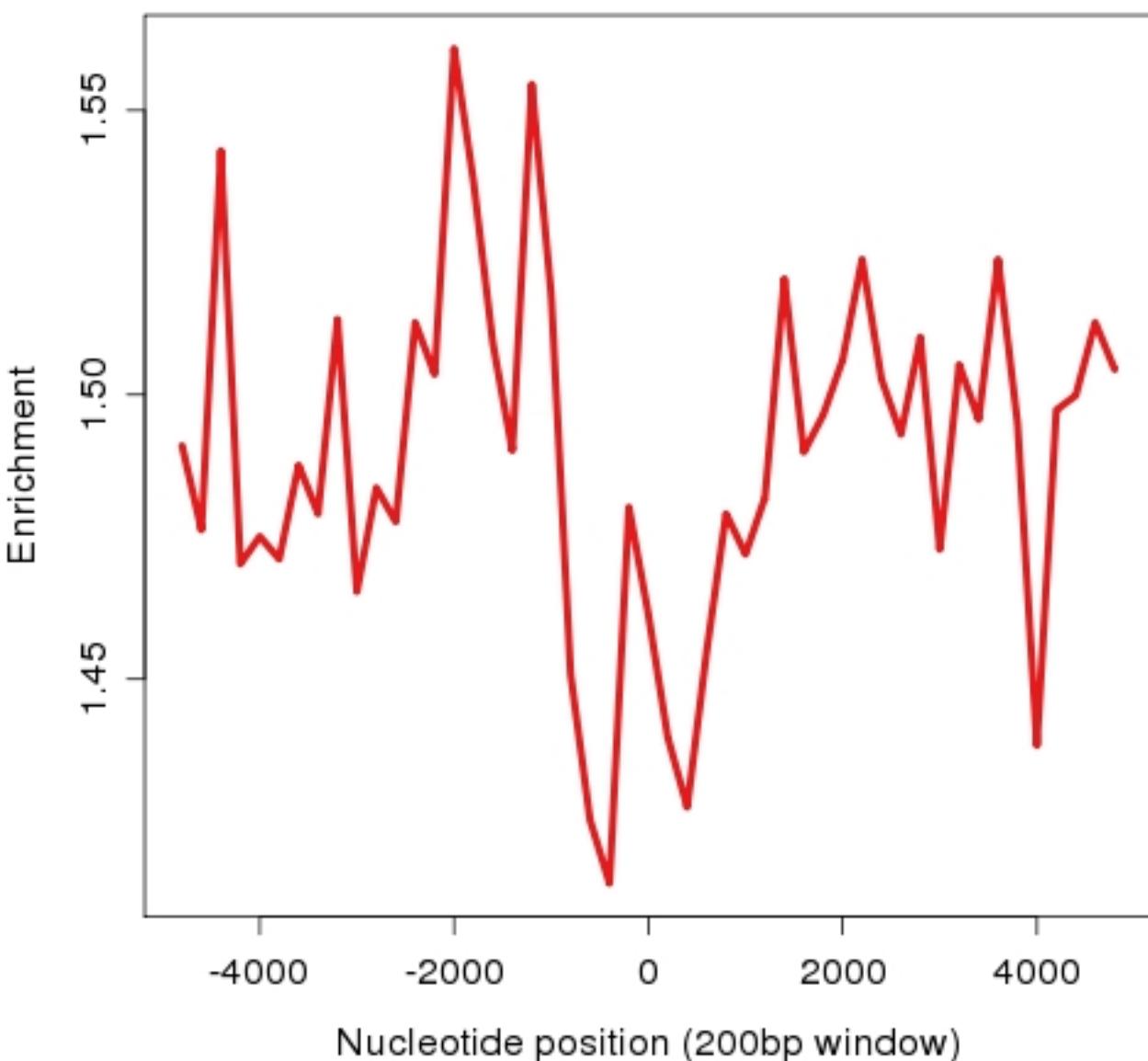


figure not available

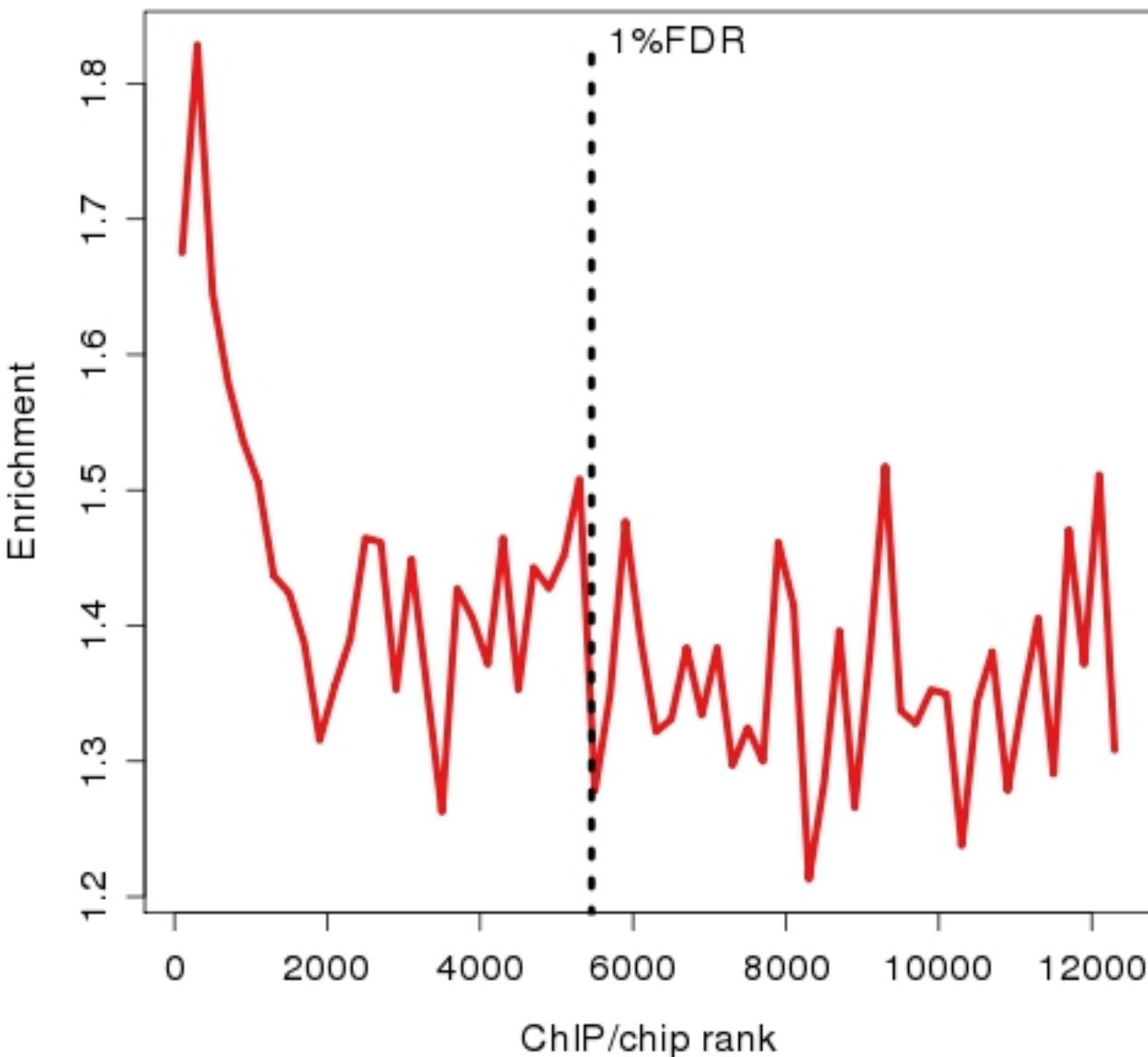
med_2 distribution of number of sites



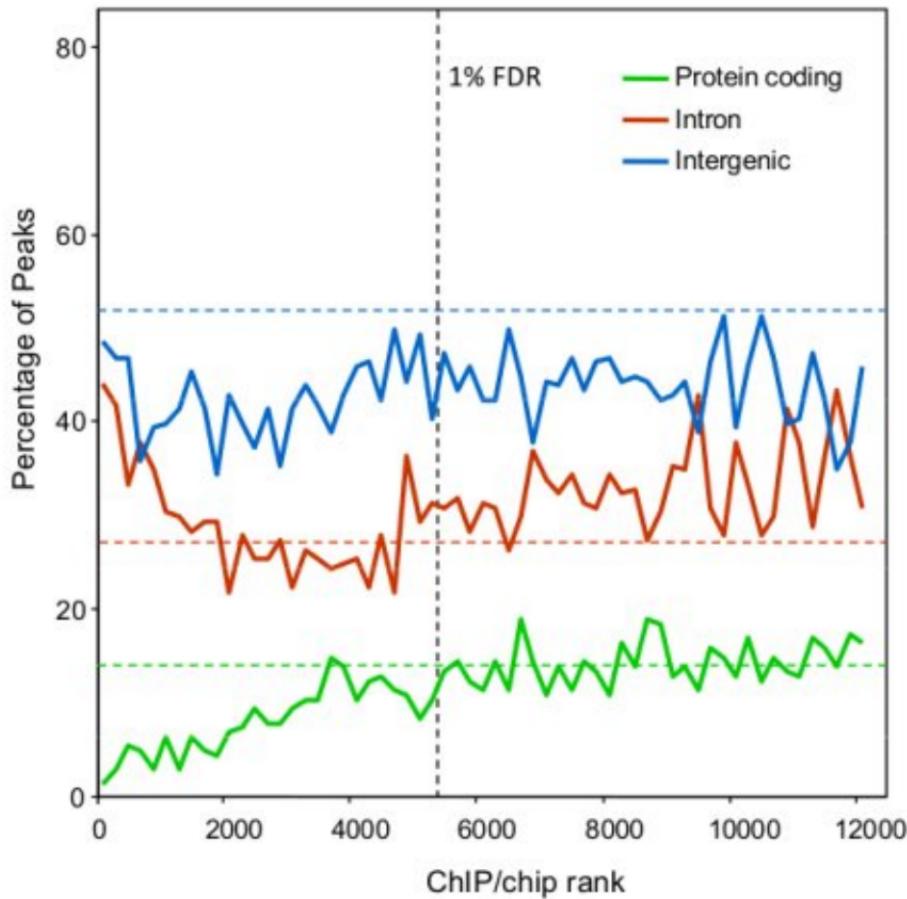
med_2 PWM enrichment across peaks



med_2 PWM enrichment down ranks



MED 2 genomic location of peaks



MED 2 GO term enrichment

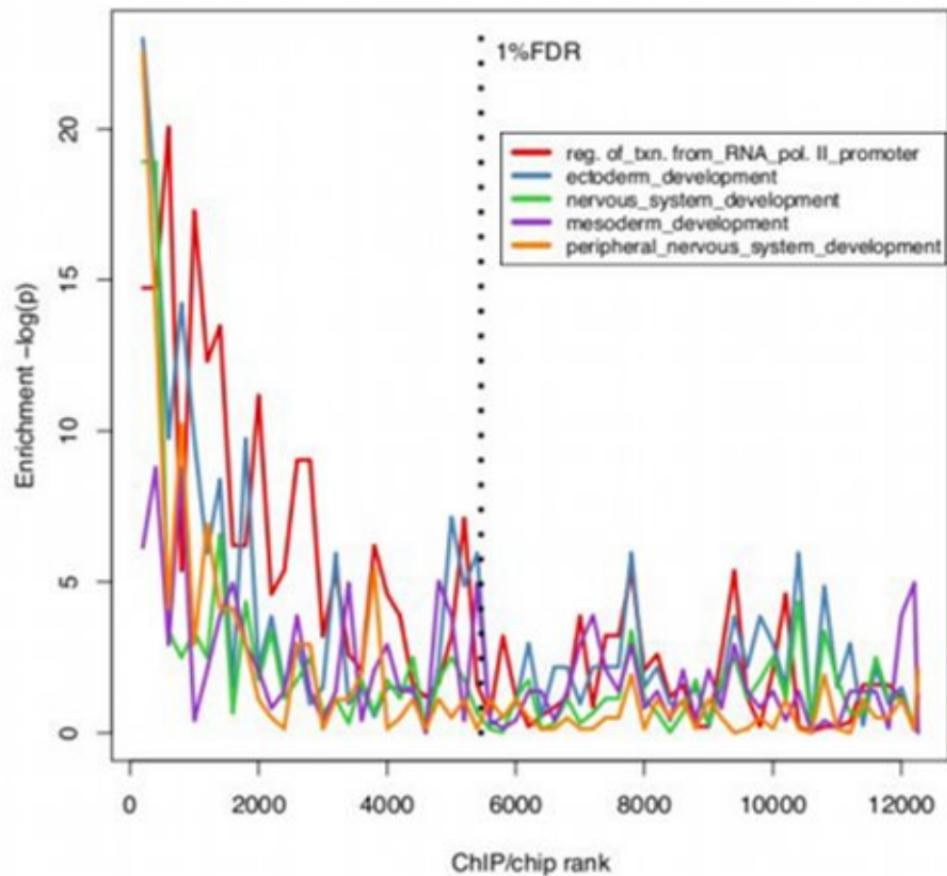
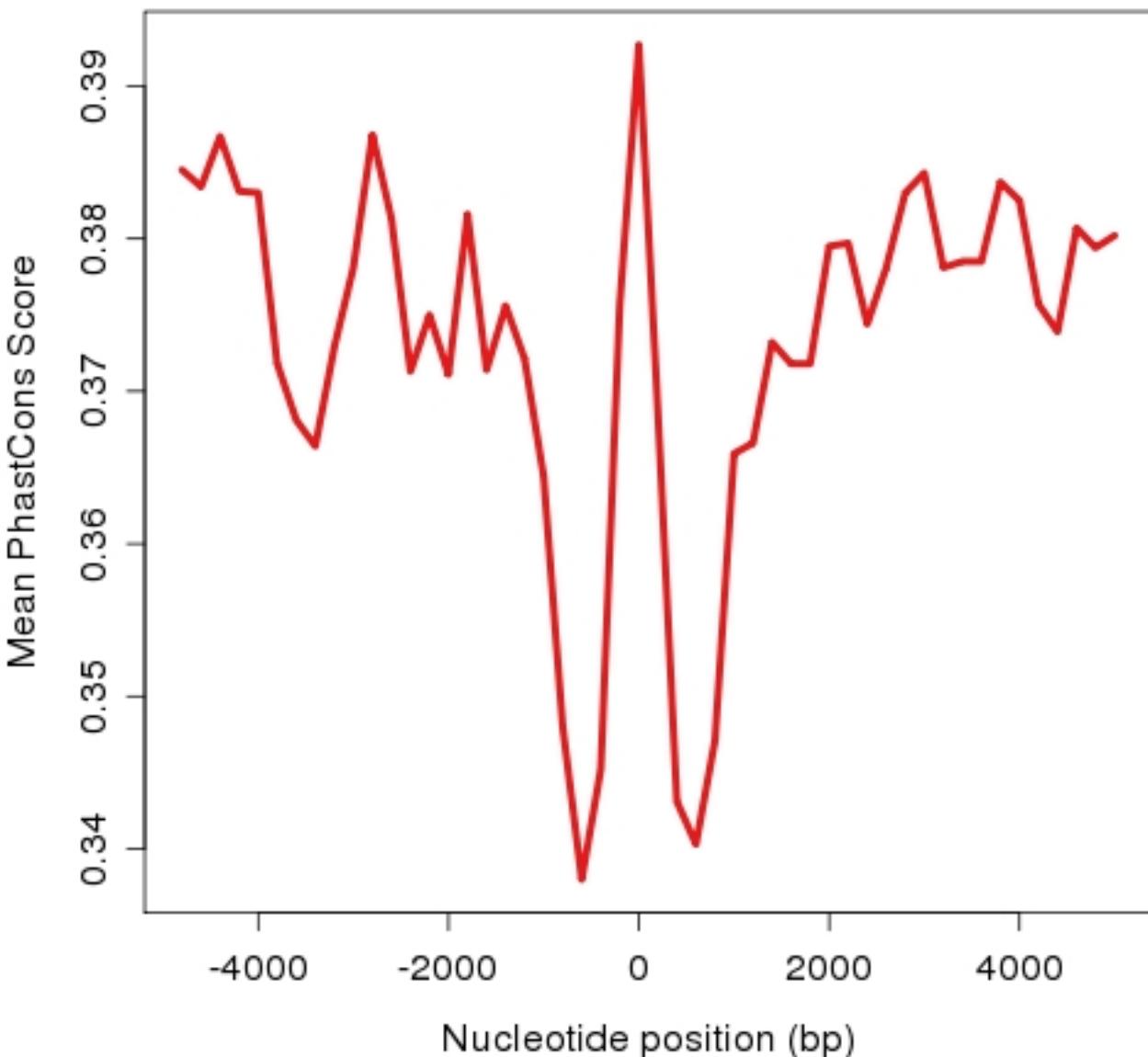


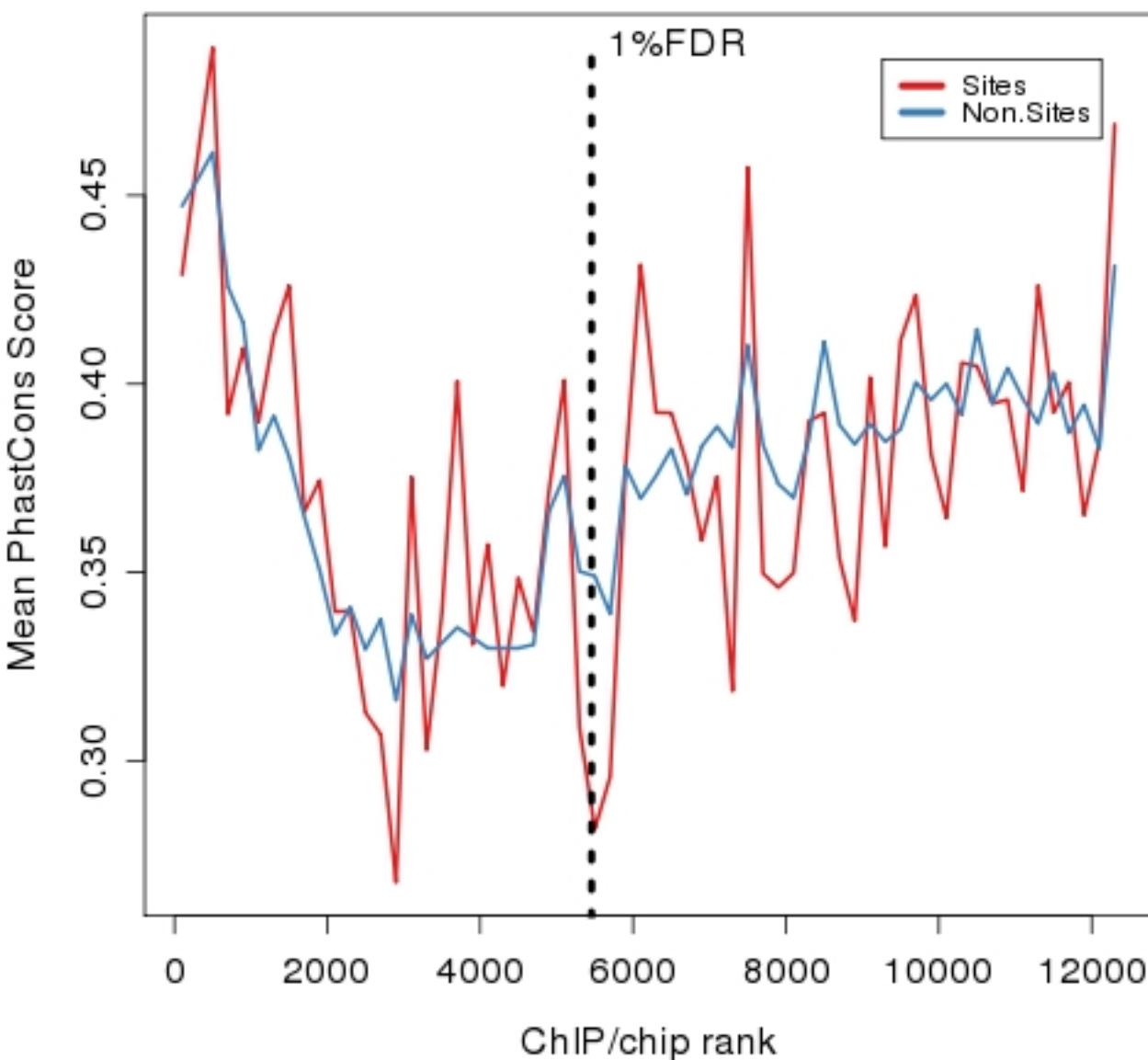
Image Not Applicable

No Matrix (PWM)
for this factor

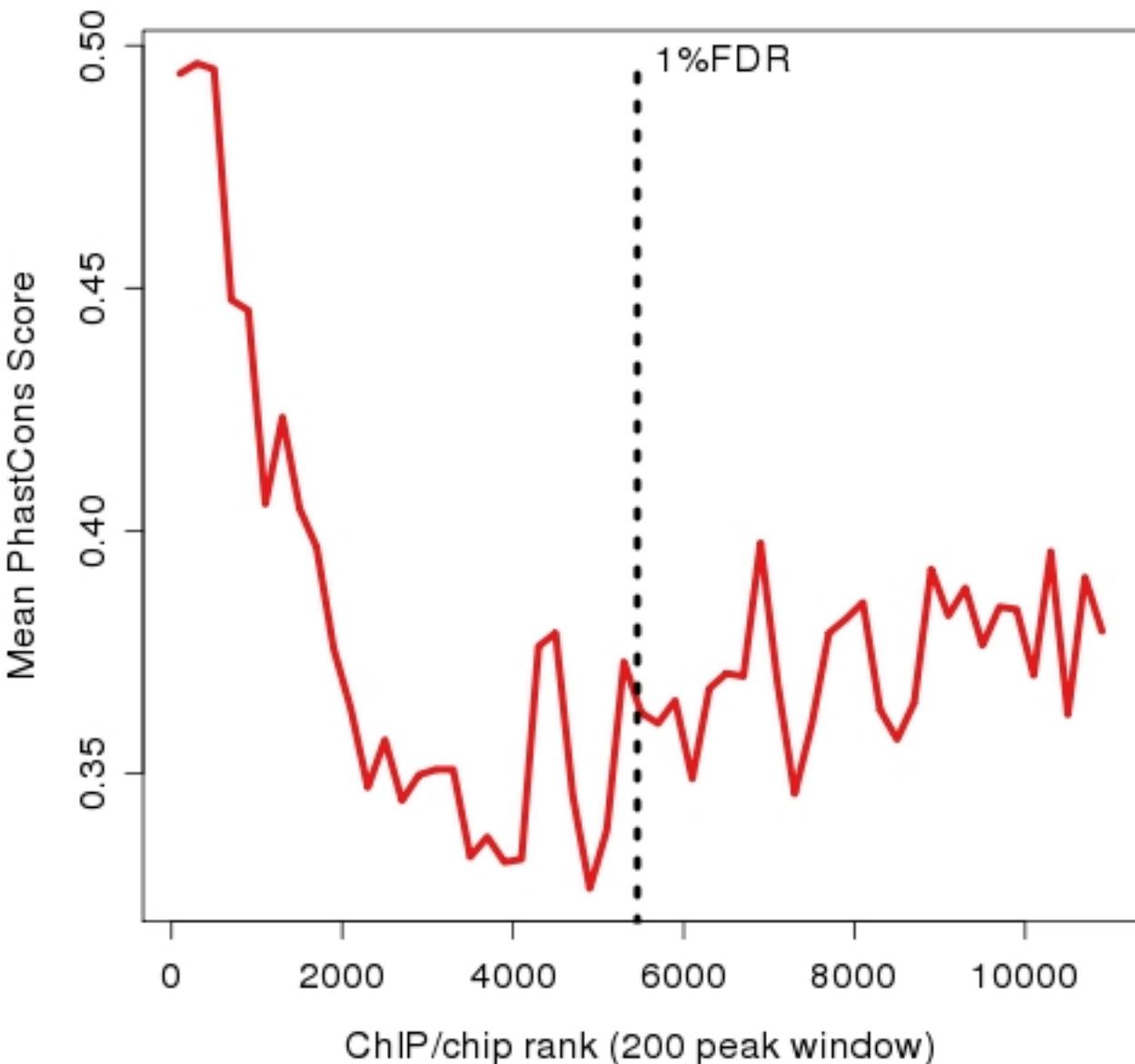
med_2 mean PhastCons across peaks



med_2 PhastCons



med_2 mean PhastCons down ranks



MED 2 GO term enrichment

